

## UGA Archaeal Collaboration Study Protocol

Thank you for participating in the first Archaeal Collaboration Study, hosted by the University of Georgia iGEM Team! Below you will find a protocol to follow for quantifying the fluorescence of your samples. Please do not hesitate to contact us should you have any questions or encounter any ambiguities.

**Important: As soon as you receive your samples in the mail, place them in a freezer (-80°C) in order to reduce potential protein damages that may incur from freezing and thawing.**

We recommend that you place your samples in the freezer and take time to read over our protocol as well as prepare your plan of action.

*You have been given 30 total samples, (i.e., triplicate cell extracts obtained from 10 colonies, and please stay in accordance with our labels throughout your experiment.)*

Each sample contains 10µL of cell extracts. The buffer used is 25mM PIPES pH 6.8. The PIPES should act as a negative control in your experiment. If PIPES is not available, a buffer with the same pH may be sufficient.

1. Remove samples from the freezer and place in a shaker at 30°C for overnight incubation. (This step is *necessary* because our cells are grown anaerobically but the fluorophore of mCherry fluorescent protein requires sufficient oxygen exposure for maturation.)
2. Using “8B” determine the linear range for your measurement device.
  - a. We recorded the highest fluorescence from sample “8B,” so if this is found to be in the linear range, then all the other samples will be as well.
  - b. Our device required a 200µL sample, so we found that a 100X dilution (i.e., 2µL cell extracts re-suspended in 198µL PIPES buffer) was within the linear range.
  - c. Determining the linear range can be done with serial dilutions (e.g., 10, 10<sup>2</sup>, 10<sup>3</sup>, 10<sup>4</sup> and 10<sup>5</sup> dilutions).
3. Our measurement device is a BioTek Synergy HT plate reader (96 well) and we use the Gen5 software.
  - a. Excitation wavelength (nm): 590/20
  - b. Emission wavelength (nm): 645/40
  - c. Scaled\* to High Wells
  - d. Setpoint Temperature: 25°C
4. Your measurement device and measurement protocol may vary from ours and other teams, so please update our form with your specific parameters. Email Steven Kodish at [skodish@uga.edu](mailto:skodish@uga.edu), Rebecca Anne Buchanan [beccab@uga.edu](mailto:beccab@uga.edu), and Akshay Chandora at [akshaych@uga.edu](mailto:akshaych@uga.edu) upon receiving this shipment for the link to the registration form and data form or for any additional questions.

\*Our values are all relative to each other (i.e., X is 20% more fluorescent than Y) so that all data can be correlated regardless of measurement device.